```
zinc fingers of the adrl gene of yeast.
CT
      Algorithms
        Amino Acid Sequence
      Bacteriophage lambda
       *DNA-Binding Proteins: GE, genetics
        Fungal Proteins: GE, genetics
      Genome
        Leucine Zippers
        Microcomputers
     Molecular Sequence Data
       *Pattern Recognition
      Protein Conformation
      Saccharomyces cerevisiae
       *Software
        User-Computer Interface
        Viral Proteins: GE, genetics
        Zinc Fingers
     0 (DNA-Binding Proteins); 0 (Fungal Proteins); 0 (Viral Proteins)
CN
GEN
    adr1
=> d his
     (FILE 'HOME' ENTERED AT 09:20:36 ON 27 JUN 2002)
                SET COST OFF
     FILE 'HCAPLUS' ENTERED AT 09:20:53 ON 27 JUN 2002
                E BRAHMACHARI S/AU
             76 S E4,E5
L1
                E DASH D/AU
             27 S E3-E7,E10
L2
                E DEBASIS/AU
                E BRAHMACHARI K/AU
              1 S E4
L3
            100 S L1-L3
L4
             10 S L4 AND MOTIF
L5
              2 S L5 AND (PROTEIN OR ?PEPTIDE?) (S) MOTIF
L6
L7
              1 S L6 NOT CLOCK/TI
     FILE 'WPIX' ENTERED AT 09:26:03 ON 27 JUN 2002
                E WO2001074130/PN
L8
              1 S E3
                E BRAHMACHARI K/AU
L9
              1 S E4
                E DASH D/AU
L10
              1 S E3
L11
              1 S L8-L10
     FILE 'HCAPLUS' ENTERED AT 09:27:21 ON 27 JUN 2002
    FILE 'WPIX' ENTERED AT 09:27:26 ON 27 JUN 2002
     FILE 'DPCI' ENTERED AT 09:27:46 ON 27 JUN 2002
                E WO2001074130/PN
     FILE 'HCAPLUS' ENTERED AT 09:28:19 ON 27 JUN 2002
              1 S ALTSCHUL ?/AU AND 1990/PY AND J MOL BIOL/JT AND (215 AND 403)
L12
              1 S CUTLER ?/AU AND 1999/PY AND MOL CELL ENDOCRINOL/JT AND (155 A
L13
              1 S GHANNOUM ?/AU AND 1999/PY AND CLIN MICROBIOL REV/JT AND (12 A
L14
              1 S MCCAFFERTY ?/AU AND 1999/PY AND CURR OPIN CHEM BIOL/JT AND (3
L15
              1 S ROOMAN ?/AU AND 1988/PY AND NATURE?/JT AND (335 AND 45)/SO
L16
              1 S WILBUR ?/AU AND 1983/PY AND PROC NATL ACAD SCI?/JT AND (80 AN
L17
              1 S WIMBERLY ?/AU AND 1999/PY AND CELL?/JT AND (97 AND 491)/SO
L18
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L19
              1 S PORSE ?/AU AND 1999/PY AND CELL?/JT AND (97 AND 423)/SO
              1 S PRESNELL ?/AU AND 1992/PY AND BIOCHEM?/JT AND (31 AND 983)/SO
L20
L21
               9 S L12-L20
     FILE 'HCAPLUS' ENTERED AT 09:48:01 ON 27 JUN 2002
     FILE 'MEDLINE' ENTERED AT 09:50:46 ON 27 JUN 2002
                E BRAHMACHARI K/AU
L22
             58 S E6-E8
                E DASH D/AU
             22 S E3-E5
L23
             77 S L22, L23
L24
              1 S L24 AND L1./CT
L25
             68 S L24 AND (D12. OR D13. OR G1. OR G5.)/CT
L26
              9 S L24 NOT L26
L27
              1 S L27 AND DNA CHIP
L28
     FILE 'MEDLINE' ENTERED AT 09:56:37 ON 27 JUN 2002
     FILE 'HCAPLUS' ENTERED AT 09:56:54 ON 27 JUN 2002
              2 S L4 AND CHIP
L29
     FILE 'HCAPLUS' ENTERED AT 09:57:17 ON 27 JUN 2002
     FILE 'BIOSIS' ENTERED AT 09:57:40 ON 27 JUN 2002
                E BRAHMACHARI/AU
L30
             79 S E9,E11-E13
                E DASH D/AU
             19 S E3-E7, E9
L31
             94 S L30, L31
L32
L33
               4 S L32 AND 00530/CC
               3 S L32 AND 04500/CC
L34
L35
               5 S L33, L34
                 SEL DN AN 1-3
              3 S L35 AND E1-E6
L36
     FILE 'BIOSIS' ENTERED AT 10:00:04 ON 27 JUN 2002
             89 S L32 NOT L35
L37
     FILE 'WPIX' ENTERED AT 10:03:12 ON 27 JUN 2002
            196 S GO6F/IC, ICM, ICS AND CO7K/IC, ICM, ICS
L38
L39
            158 S L38 AND (B12-K04? OR C12-K04?)/MC
              1 S L38 AND T01-E01B/MC
L40
              1 S CO7K/IC, ICM, ICS AND T01-E01B/MC
L41
             50 S CO7K/IC, ICM, ICS AND T01-J05B?/MC
L42
                 SEL DN AN 9 10 28 20 41 45 48 50 L42
              8 S L42 AND E7-E29
L43
L44
            124 S L39 NOT L40-L43
                 SEL DN AN 13 2 31 34 37 61 97 122
              8 S L44 AND E30-E51
L45
L46
             16 S L43, L45
L47
             32 S L38 NOT L39-L46
                 SEL DN AN 20
L48
               1 S L47 AND E52-E45
L49
             17 S L40, L41, L46, L48
L50
             16 S L49 NOT L11
             60 S GO6F/IC, ICM, ICS AND MOTIF
L51
L52
              4 S GO6F/IC, ICM, ICS, ICA, ICI AND (M OR MYCO?)()TUBERCUL?
L53
              3 S L52 NOT L11
L54
              0 S L51 AND L52 NOT L11
L55
             19 S L50, L53
            175 S (T01-? OR T02-? OR T03-?)/MC AND C07K/IC, ICM, ICS
L56
             44 S L56 NOT L38-L55
L57
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SEL DN AN 29
              1 S L57 AND E55-E57
L58
L59
             20 S L55, L58 AND L38-L58
             18 S L59 AND (MOTIF OR ?OVERLAP? OR LENGTH OR INVAR? OR CONSERV? O
L60 .
             20 S L59, L60 NOT L11
L61
     FILE 'WPIX' ENTERED AT 10:52:21 ON 27 JUN 2002
     FILE 'BIOSIS' ENTERED AT 10:52:43 ON 27 JUN 2002
          18750 S (M OR MYCO?) () TUBERCUL?
L62
          57156 S MYCOBACTERIACEAE+NT/BC
L63
L64
          60129 S L62, L63
            514 S L64 AND 00530/CC
L65
             35 S L65 AND (10054 OR 10064 OR 13012)/CC
                SEL DN AN 2 16
L67
              2 S E58-E61 AND L66
            139 S L65 AND *00530/CC NOT L66
                SEL DN AN 2 4 9 12 34
L69
              5 S L68 AND E62-E71
L70
              7 S L67, L69
     FILE 'BIOSIS' ENTERED AT 11:02:57 ON 27 JUN 2002
     FILE 'MEDLINE' ENTERED AT 11:03:20 ON 27 JUN 2002
          20194 S L62
                E MYCOBACTERIUM TUBERCULOSIS/CT
                E E3+ALL
          13804 S E14+NT
L73
          20194 S L71, L72
            167 S L1.700./CT AND L73
L74
             11 S L1.725./CT AND L73
L75
            168 S L74, L75
L76
            118 S L73 AND MOTIF
L77
L78
              4 S L76 AND L77
L79
             67 S L77 AND AMINO ACID SEQUENCE+NT/CT
                E AMINO ACID MOTIF/CT
                E E4+ALL
          11359 S E38+NT
L80
L81
          17706 S E63+NT
           6425 S E62+NT
L82
L83
          29020 S E37+NT
           3469 S (L1.700. OR L1.725.)/CT AND L80-L83
L84
         268504 S L1.700./CT OR L1.725./CT
L85
            835 S L84 AND L85/MAJ
L86
            299 S L86 AND (L80/MAJ OR L81/MAJ OR L82/MAJ OR L83/MAJ)
L87
             32 S L87 NOT AB/FA
L88
                E SEQUENCE HOMOLOGY/CT
                E E6+ALL
          56315 S E47+NT
L89
L90
             43 S L87 AND L89
            164 S L86 AND MOTIF
L91
             56 S L91 AND (CONSERV? OR INVAR?)
L92
L93
              4 S L92 NOT AB/FA
L94
             52 S L92 NOT L93
             45 S L94 AND (AMINO ACID SEQUENCE+NT OR D12.)/CT
L95
              7 S L94 NOT L95
L96
                SEL DN AN 2 6
              2 S E1-E6
L97
L98
             52 S L94, L97
                SEL DN AN 1 2 4 7 13 20 22 25 26 28 29 35 36 37 40 47
L99
             36 S L98 NOT E7-E54
L100
             50 S L91 AND L87
             35 S L100 NOT L98
L101
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SEL DN AN 3 4 6 8 10 12 14-18 20-22 30-35 20 S L101 AND E55-E114

L102 L103

56 S L99,L102 56 S L103 AND L71-L103 L104